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## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/037,311A

DATE: 12/10/2002 P.6

TIME: 15:43:38

Input Set : A:\MS00-001C2 XFTASE.txt

Output Set: N:\CRF4\12102002\J037311A.raw

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3 <110> APPLICANT: MICHIGAN STATE UNIVERSITY
5 <120> TITLE OF INVENTION: XYLOGLUCAN FUCOSYLTRANSFERASES
7 <130> FILE REFERENCE: MS00-001C2
9 <140> CURRENT APPLICATION NUMBER: US 10/037,311A
10 <141> CURRENT FILING DATE: 2001-11-09
12 <150> PRIOR APPLICATION NUMBER: US60/117,555
13 <151> PRIOR FILING DATE: 1999-01-28
15 <160> NUMBER OF SEQ ID NOS: 15
17 <170> SOFTWARE: PatentIn version 3.1
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20 <211> LENGTH: 558
21 <212> TYPE: PRT
22 <213> ORGANISM: Arabidopsis thaliana
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31 20 25 30
34 Tyr Leu Ser Ser Gly Thr Met Lys Leu Thr Arg Thr Phe Thr Thr Cys
35 35 40 45
38 Leu Ile Val Phe Ser Val Leu Val Ala Phe Ser Met Ile Phe His Gln
39 50 55 60
42 His Pro Ser Asp Ser Asn Arg Ile Met Gly Phe Ala Glu Ala Arg Val
43 65 70 75 80
46 Leu Asp Ala Gly Val Phe Pro Asn Val Thr Asn Ile Asn Ser Asp Lys
47 85 90 95
50 Leu Leu Gly Gly Leu Leu Ala Ser Gly Phe Asp Glu Asp Ser Cys Leu
51 100 105 110
54 Ser Arg Tyr Gln Ser Val His Tyr Arg Lys Pro Ser Pro Tyr Lys Pro
55 115 120 125
58 Ser Ser Tyr Leu Ile Ser Lys Leu Arg Asn Tyr Glu Lys Leu His Lys
59 130 135 140
62 Arg Cys Gly Pro Gly Thr Glu Ser Tyr Lys Lys Ala Leu Lys Gln Leu
63 145 150 155 160
66 Asp Gln Glu His Ile Asp Gly Asp Gly Glu Cys Lys Tyr Val Val Trp
67 165 170 175
70 Ile Ser Phe Ser Gly Leu Gly Asn Arg Ile Leu Ser Leu Ala Ser Val
71 180 185 190
74 Phe Leu Tyr Ala Leu Leu Thr Asp Arg Val Leu Leu Val Asp Arg Gly
75 195 200 205
78 Lys Asp Met Asp Asp Leu Phe Cys Glu Pro Phe Leu Gly Met Ser Trp
79 210 215 220
82 Leu Leu Pro Leu Asp Phe Pro Met Thr Asp Gln Phe Asp Gly Leu Asn

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83 225          230          235          240
86 Gln Glu Ser Ser Arg Cys Tyr Gly Tyr Met Val Lys Asn Gln Val Ile
87          245          250          255
90 Asp Thr Glu Gly Thr Leu Ser His Leu Tyr Leu His Leu Val His Asp
91          260          265          270
94 Tyr Gly Asp His Asp Lys Met Phe Phe Cys Glu Gly Asp Gln Thr Phe
95          275          280          285
98 Ile Gly Lys Val Pro Trp Leu Ile Val Lys Thr Asp Asn Tyr Phe Val
99          290          295          300
102 Pro Ser Leu Trp Leu Ile Pro Gly Phe Asp Asp Glu Leu Asn Lys Leu
103 305          310          315          320
106 Phe Pro Gln Lys Ala Thr Val Phe His His Leu Gly Arg Tyr Leu Phe
107          325          330          335
110 His Pro Thr Asn Gln Val Trp Gly Leu Val Thr Arg Tyr Tyr Glu Ala
111          340          345          350
114 Tyr Leu Ser His Ala Asp Glu Lys Ile Gly Ile Gln Val Arg Val Phe
115          355          360          365
118 Asp Glu Asp Pro Gly Pro Phe Gln His Val Met Asp Gln Ile Ser Ser
119          370          375          380
122 Cys Thr Gln Lys Glu Lys Leu Leu Pro Glu Val Asp Thr Leu Val Glu
123 385          390          395          400
126 Arg Ser Arg His Val Asn Thr Pro Lys His Lys Ala Val Leu Val Thr
127          405          410          415
130 Ser Leu Asn Ala Gly Tyr Ala Glu Asn Leu Lys Ser Met Tyr Trp Glu
131          420          425          430
134 Tyr Pro Thr Ser Thr Gly Glu Ile Ile Gly Val His Gln Pro Ser Gln
135          435          440          445
138 Glu Gly Tyr Gln Gln Thr Glu Lys Lys Met His Asn Gly Lys Ala Leu
139          450          455          460
142 Ala Glu Met Tyr Leu Leu Ser Leu Thr Asp Asn Leu Val Thr Ser Ala
143 465          470          475          480
146 Trp Ser Thr Phe Gly Tyr Val Ala Gln Gly Leu Gly Gly Leu Lys Pro
147          485          490          495
150 Trp Ile Leu Tyr Arg Pro Glu Asn Arg Thr Thr Pro Asp Pro Ser Cys
151          500          505          510
154 Gly Arg Ala Met Ser Met Glu Pro Cys Phe His Ser Pro Pro Phe Tyr
155          515          520          525
158 Asp Cys Lys Ala Lys Thr Gly Ile Asp Thr Gly Thr Leu Val Pro His
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163 545          550          555
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168 <212> TYPE: DNA
169 <213> ORGANISM: Arabidopsis thaliana
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174 aagtccgtta atttctccga actacttcaa atgaagtatc tcagctccgg tacgatgaag    120
176 ctcacgagaa ccttcactac ttgcttgata gtcttctctg tactagtagc attctcaatg    180

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180 ctcgacgccg gagttttccc aaattctgat aagcttctcg gagggctact tgcttctggg 300
182 tttgatgaag attcttgccct tagtaggtac caatcagttc attaccgtaa accttcacct 360
184 tacaagccat ctctctatct catctctaag cttagaaact acgaaaagct tcacaagcga 420
186 tgtgggtccg gtactgaatc ttacaagaaa gctctaaaac aacttgatca agaacatatt 480
188 gatgggtgat gtgaatgcaa atatggtgtg tggatttctt ttagcggctt agggaaacagg 540
190 atactttctc tagcctcggg ttttctttac gcgcttttaa cggatagagt cttgcttggt 600
192 gaccgaggga aagacatgga tgatctcttt tgcgagccgt ttctcgggat gtcgtggttg 660
194 ctacctttag atttccctat gactgatcag tttgatggat taaatcaaga atcatctcgt 720
196 tgttatggat atatggtgaa gaatcagggt attgatactg agggaaacttt gtctcatctt 780
198 tatcttcac tttgttcata ttatggagat catgataaga tgttcttctg tgaaggagac 840
200 caaacattca tcgggaaagt cccttgggtt attgttaaaa cagacaatta ctttgttcca 900
202 tctctgtggt taataccggg tttcgatgat gaactaaaca agctattccc acagaaagcg 960
204 actgtctttc atcacttagg taggtatctt tttcacccaa ctaaccaagt atggggctta 1020
206 gtcactagat actacgaagc ttacttatcg catgcggatg agaagattgg gattcaagta 1080
208 agagttttcg atgaagaccc ggggtccattt cagcatgtga tggatcagat ttcactttgt 1140
210 actcaaaaag agaaacttct acctgaagta gacacactag tggagagatc tcgccatgtt 1200
212 aataccccc aacacaaagc cgtgcttgtc acatctttga acgcgggtta cgcggagAAC 1260
214 ttaaagagta tgtattggga atatccgaca tcaactggag aaatcatcgg tgttcacag 1320
216 ccgagccaag aaggttatca gcagaccgaa aaaaagatgc ataatggcaa agctcttgcg 1380
218 gaaatgtatc ttttgagttt gacagataat cttgtgacaa gtgcttggtc tacatttggg 1440
220 tatgtagctc aagggtcttg aggttttaaag ccttggtatc tctatagacc cgaAACaccgt 1500
222 acaactccc atccttcgtg tggtcgggct atgtcgatgg agccttgttt ccactcgcct 1560
224 ccattctatg attgtaaagc gaaaacgggt attgacacgg gaacactagt tcctcatgtg 1620
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232 <213> ORGANISM: Arabidopsis thaliana
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237 aagtccgtta atttctccga actacttcaa atgaagtatc tcagctccgg tacgatgaag 120
239 ctcacgagaa ctttactac ttgcttgata gtcttctctg tactagtagc attctcaatg 180
241 atctttcacc aacacccatc tgattcaaat cggattatgg gtttcgccga agctagagtt 240
243 ctcgacgccg gagttttccc aaatgttact aacatcagta tgtgttcttc caagtcaaag 300
245 ttttgagctt tattacttta gatctcgttc tttactactac gcatttgcct ctgtatgtcc 360
247 atagctcttg gtcgatttca atttgagatc tatactcata aaaattgagt ctttgtcagt 420
249 cacaagacta ctatttttgg tttgatgttg ttttggtgaa aaagtgtctt tttgttttgg 480
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253 ttttgttttg tagattctga taagcttctc ggagggtctac ttgcttctgg ttttgatgaa 600
255 gattcttgcc ttagtaggta ccaatcagtt cattaccgta aaccttcacc ttacaagcca 660
257 tcttcttata tcatctctaa gcttagaaac tacgaaaagc ttcacaagcg atgtgggtccg 720
259 ggtactgaat cttacaagaa agctctaaaa caacttgatc aagaacatat tgatgggtgat 780
261 ggtgaatgca aatatgttgt gtggatttct tttagcggct tagggaaacag gatactttct 840
263 ctagcctcgg ttttctttaa cgcgctttta acggatagag tcttgccttg tgaccgaggg 900
265 aaagacattg atgatctctt ttgcgagccg tttctcggta tgtcgtggtt gctaccttta 960
267 gatttcccta tgactgatca gtttgatgga ttaaatcaag aatcatctcg ttgttatgga 1020
269 tatatgggtg agaatacagg gattgatact gagggaaact tgtctcatct ttatcttcac 1080
271 cttgttcacg attatggaga tcatgataag atgttcttct gtgaaggaga ccaaacattc 1140

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DATE: 12/10/2002

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Output Set: N:\CRF4\12102002\J037311A.raw

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277 catcacttag gtaggtatct ttttcaccca actaaccaag tatggggctt agtcactaga 1320
279 tactacgaag cttacttatc gcatgcggat gagaagattg ggattcaagt aagagttttc 1380
281 gatgaagacc cgggtccatt tcagcatgtg atggatcaga tttcatcttg tactcaaaaa 1440
283 gagaaacttc tacctgaagt agacacacta gtggagagat ctgccatgt taataccccc 1500
285 aaacacaaag ccgtgcttgt cacatctttg aacgcggggt acgcggagaa cttaaagagt 1560
287 atgtattggg aatatccgac atcaactgga gaaatcatcg gtgttcatca gccgagccaa 1620
289 gaaggttatc agcagaccga aaaaaagatg cataatggca aagctcttgc ggaaatgtat 1680
291 cttttgagtt tgacagataa tcttgtgaca agtgcttggg ctacatttgg atatgtagct 1740
293 caaggtcttg gaggtttaaa gccttggata ctctatagac ccgaaaaccg tacaactccc 1800
295 gatccttcgt gtggtcgggc tatgtcgatg gaggcttgtt tccactcgcc tccattctat 1860
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312 atggaaccaaa atgtgacatc ctcatcagat gattcatcac tagtgacagag agatcaagaa 180
314 caaaaaaggta aacttacttt cttctttttt ttttgaaatg tttctaaatt tttctttgaa 240
316 tgtttcatca gattctgtag atatgtctct gcttggaggg ctacttgtat ctggtttcaa 300
318 gaaaagagtct tgcttgagta gataccaatc ttacctctac cgtaaagctt caccgtataa 360
320 accttcgttg catctacttt cgaagcttag agcttacgaa gagcttcata aaagatgtgg 420
322 accgggaaca agacagtata ccaatgcaga aagattgctt aaacagaaac aaacagggtga 480
324 gatggaatca caaggatgca agtatgttgt ttggatgtcg tttagcggat taggaaacag 540
326 gattatcagt attgcttctg tgtttctgta tgcaatgttg acagatagag tcttgcttgt 600
328 tgaaggaggg gaacagttcg cggtatttatt ctgcgaaccg ttcctcgata ccacttggtt 660
330 actaccgaaa gatttcacct tagctagtca gttcagtggc tttggtcaaa actcagctca 720
332 ctgccatgga gatatgctga agaggaaaact gattaatgaa tcctctgttt cgtctctgtc 780
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336 agaagatcaa aatctcttaa agaattgtcc ttggttgatc atgaggacaa acaacttctt 900
338 tgaccagtct cttttcttga tttcttcttt cgaagaagag ctcggtatga tgtttcccga 960
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342 gggactaatc acaagatact atcaagctta cttagccaaa gctgatgaaa ggattgggtct 1080
344 tcaaataaga gtctttgatg agaaatccgg cgtatctcct cgagtcacaa agcaaatacat 1140
346 ttcgtgtgtt caaaacgaga atctgttacc gagactaagc aaaggtgaag aacaatacaa 1200
348 gcagccatca gaagaagagt tgaaactcaa atctgtcttg gtcacctctt taacaacagg 1260
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## RAW SEQUENCE LISTING

DATE: 12/10/2002

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TIME: 15:43:38

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Output Set: N:\CRF4\12102002\J037311A.raw

368 &lt;211&gt; LENGTH: 538

369 &lt;212&gt; TYPE: PRT

370 &lt;213&gt; ORGANISM: Arabidopsis thaliana

372 &lt;400&gt; SEQUENCE: 5

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382 Val Gln Ala Ser Arg Phe Ile Thr Met Glu Pro Asn Val Thr Ser Ser
383          35          40          45
386 Ser Asp Asp Ser Ser Leu Val Gln Arg Asp Gln Glu Lys Asp Ser
387          50          55          60
390 Val Asp Met Ser Leu Leu Gly Gly Leu Leu Val Ser Gly Phe Lys Lys
391 65          70          75          80
394 Glu Ser Cys Leu Ser Arg Tyr Gln Ser Tyr Leu Tyr Arg Lys Ala Ser
395          85          90          95
398 Pro Tyr Lys Pro Ser Leu Leu Leu Ser Lys Leu Arg Ala Tyr Glu Glu
399          100         105         110
402 Leu His Lys Arg Cys Gly Pro Gly Thr Arg Gln Tyr Thr Asn Ala Glu
403          115         120         125
406 Arg Leu Leu Lys Gln Lys Gln Thr Gly Glu Met Glu Ser Gln Gly Cys
407          130         135         140
410 Lys Tyr Val Val Trp Met Ser Phe Ser Gly Leu Gly Asn Arg Ile Ile
411 145         150         155         160
414 Ser Ile Ala Ser Val Phe Leu Tyr Ala Met Leu Thr Asp Arg Val Leu
415         165         170         175
418 Leu Val Glu Gly Gly Gln Phe Ala Asp Leu Phe Cys Glu Pro Phe
419         180         185         190
422 Leu Asp Thr Thr Trp Leu Leu Pro Lys Asp Phe Thr Leu Ala Ser Gln
423         195         200         205
426 Phe Ser Gly Phe Gly Gln Asn Ser Ala His Cys His Gly Asp Met Leu
427         210         215         220
430 Lys Arg Lys Leu Ile Asn Glu Ser Ser Val Ser Ser Leu Ser His Leu
431 225         230         235         240
434 Tyr Leu His Leu Ala His Asp Tyr Asn Glu His Asp Lys Met Phe Phe
435         245         250         255
438 Cys Glu Glu Asp Gln Asn Leu Leu Lys Asn Val Pro Trp Leu Ile Met
439         260         265         270
442 Arg Thr Asn Asn Phe Phe Ala Pro Ser Leu Phe Leu Ile Ser Ser Phe
443         275         280         285
446 Glu Glu Glu Leu Gly Met Met Phe Pro Glu Lys Gly Thr Val Phe His
447         290         295         300
450 His Leu Gly Arg Tyr Leu Phe His Pro Ser Asn Gln Val Trp Gly Leu
451 305         310         315         320
454 Ile Thr Arg Tyr Tyr Gln Ala Tyr Leu Ala Lys Ala Asp Glu Arg Ile
455         325         330         335
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459         340         345         350
462 Val Thr Lys Gln Ile Ile Ser Cys Val Gln Asn Glu Asn Leu Leu Pro

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RAW SEQUENCE LISTING ERROR SUMMARY      DATE: 12/10/2002  
PATENT APPLICATION: US/10/037,311A      TIME: 15:43:39

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Output Set: N:\CRF4\12102002\J037311A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:6; N Pos. 10,29,61,92,101,133,147,168,197,215  
Seq#:7; Xaa Pos. 10,20,31,34,49,56,66,72  
Seq#:8; N Pos. 4,29,146,190,195,224,263,354,382,383,397,408,426,433,434,438  
Seq#:8; N Pos. 454,481,489,511  
Seq#:11; N Pos. 148,150,221,248,330,382,410,422,451,502,509,528,539,549,647  
Seq#:11; N Pos. 650,659,701,702  
Seq#:12; N Pos. 276,361,386,409,433,481,490  
Seq#:15; N Pos. 4,5,12,16,27,50,53,72,73,81,87,98,118,147,153,159,168,205  
Seq#:15; N Pos. 275

## VERIFICATION SUMMARY

DATE: 12/10/2002

PATENT APPLICATION: US/10/037,311A

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Output Set: N:\CRF4\12102002\J037311A.raw

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L:578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:60  
L:580 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:120  
L:582 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:180  
L:642 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0  
L:646 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:16  
L:650 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:32  
L:654 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:48  
L:658 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:64  
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L:784 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:120  
L:786 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:180  
L:788 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:240  
L:790 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:300  
L:792 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:360  
L:794 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:420  
L:796 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:480  
L:979 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:120  
L:981 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:180  
L:983 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:240  
L:985 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:300  
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L:1068 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:480  
L:1211 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0  
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L:1215 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:120  
L:1217 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:180  
L:1219 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:240